

1600

RAW SEQUENCE LISTING

26 <170> SOFTWARE: PatentIn version 3.0

DATE: 10/17/2003

PATENT APPLICATION: US/09/759,152A

TIME: 16:43:36

Input Set : A:\PTO.YF.txt

Output Set: N:\CRF4\10172003\I759152A.raw

```
4 <110> APPLICANT: Palmer, Michelle A.J.
         Gee, Melissa
         Tillotson, Bonnie
 6
        Chang, Xiao-Jia
 9 <120> TITLE OF INVENTION: Systems for Sensitive Detection of G-Protein Coupled
         Receptor and Orphan Receptor Function Using Reporter
10
         Enzyme Mutant Complementation
11
13 <130> FILE REFERENCE: 4085-235-27 CIP
15 <140> CURRENT APPLICATION NUMBER: US 09/759,152A
16 <141> CURRENT FILING DATE: 2001-01-16
18 <150> PRIOR APPLICATION NUMBER: US 09/654,499
19 <151> PRIOR FILING DATE: 2000-09-01
21 <150> PRIOR APPLICATION NUMBER: US 60/180,669
22 <151> PRIOR FILING DATE: 2000-02-07
24 <160> NUMBER OF SEQ ID NOS: 10
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2685 <210> SEQ ID NO: 10 2686 <211> LENGTH: 5 212> PRT 2687 <213> ORGANISM: Artificial Sequence W--> 2689 <220> FEATURE: <223> GS Linker E--> 2691 <212> TYPE: hard return W--> 2691 <223> OTHER INFORMATION: W--> 2691 <400> 10 10 10 Gly Gly Gly Ser, 1 5 2692 4085-235-27CIP.ST25 2696 Page 1

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/09/759,152A

DATE: 10/17/2003 TIME: 16:43:37

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Use of <220> Feature (NEW RULES):

Sequence(s) are missing the <220> Feature and associated headings.

Use of <220> to <223> is MANDATORY if <213> ORGANISM is "Artificial Sequence" or "Unknown". Please explain source of genetic material in <220> to <223> section (See "Federal Register," 6/01/98, Vol. 63, No. 104,pp.29631-32) (Sec.1.823 of new Rules)

Seq#:10

VERIFICATION SUMMARY

DATE: 10/17/2003 PATENT APPLICATION: US/09/759,152A TIME: 16:43:37

Input Set : A:\PTO.YF.txt

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L:2112 M:283 W: Missing Blank Line separator, <220> field identifier L:2689 M:256 W: Invalid Numeric Header Field, <220> has non-blank data L:2691 M:282 E: Numeric Field Identifier Missing, <212> is required.

L:2691 M:258 W: Mandatory Feature missing, <223> Tag not found for SEQ#:10, <213>

ORGANISM: Artificial Sequence

L:2691 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:10, Line#:2691